



(12) **EUROPEAN PATENT APPLICATION**

(43) Date of publication:  
**29.12.2004 Bulletin 2004/53**

(51) Int Cl.<sup>7</sup>: **C07H 7/027, C07H 1/00**

(21) Application number: **03013457.1**

(22) Date of filing: **24.06.2003**

(84) Designated Contracting States:  
**AT BE BG CH CY CZ DE DK EE ES FI FR GB GR**  
**HU IE IT LI LU MC NL PT RO SE SI SK TR**  
Designated Extension States:  
**AL LT LV MK**

• **Marliere, Philippe**  
**91450 Etolles (FR)**

(72) Inventor: **The designation of the inventor has not yet been filed**

(71) Applicants:  
• **Evologic S.A.**  
**91000 Evry (FR)**

(74) Representative: **VOSSIUS & PARTNER**  
**Siebertstrasse 4**  
**81675 München (DE)**

(54) **Production of 2'-deoxynucleosides and 2'-deoxynucleoside precursors from 2-dehydro-3-deoxy-D-gluconate**

(57) This invention relates to a process for preparing 2'-deoxynucleoside compounds or 2'-deoxynucleoside precursors using 2-dehydro-3-deoxy-D-gluconic acid (usually abbreviated as KDG) or its salts as a starting material. A variety of 2'-deoxynucleosides and their an-

alogues are used as a starting material for synthesis or drug formulation in production of an antiviral, anticancer or antisense agent.

## Description

**[0001]** This invention relates to a process for preparing 2'-deoxynucleoside compounds or 2'-deoxynucleoside precursors using 2-dehydro-3-deoxy-D-gluconic acid (usually abbreviated as KDG) or its salts as a starting material. A variety of 2'-deoxynucleosides and their analogues are used as a starting material for synthesis or drug formulation in production of an antiviral, anticancer or antisense agent.

**[0002]** Specifically, the invention relates to a method in which KDG or a derivative of KDG is subjected to a decarboxylation step to remove the original carboxy group of KDG. In a preferred embodiment, the KDG used in the method according to the invention is enzymatically produced from D-gluconate or D-glucosamine.

**[0003]** 2'-deoxynucleosides and 2'-deoxynucleoside precursors including 2-deoxy-D-ribose are used as starting material for synthesis or drug formulation, for instance, in production of antiviral and anticancer agent. 2'-deoxynucleosides or derivatives thereof and 2'-deoxynucleoside precursors are also used as reagents for research, diagnosis and synthesis of therapeutic antisense molecules.

**[0004]** In one method of the prior art, deoxynucleosides are generated from biological materials such as testis (WO 99/49074) or yeast or fish sperm by enzymatic cleavage of DNA. This method, however, involves several disadvantages, in particular regarding difficulties of obtaining the starting material in sufficient quantity and quality.

**[0005]** The main production process of 2-deoxy-D-ribose currently consists in chemical hydrolysis of DNA. In this case, the deoxyribosyl moiety originates in ribonucleotide reductase activity. No synthesis of 2-deoxy-D-ribose from KDG has been yet described.

**[0006]** In most living cells, deoxyribonucleosides result from a "salvage pathway" of the nucleotide metabolism. The deoxyribose moiety of deoxyribonucleosides is obtained through the reduction of a ribosyl moiety into di- or triphosphate ribonucleotides catalyzed by ribonucleotide reductases. However, the deoxyribose moiety is not recycled, but is degraded into D-glyceraldehyde-3-phosphate and acetaldehyde following the reactions of central metabolism:

- deoxynucleoside is cleaved into deoxyribose-1-phosphate and nucleobase through phosphorolysis mediated by products of the genes encoding thymidine phosphorylase (deoA), purine-nucleoside phosphorylase (deoD), uridine phosphorylase (udp) or xanthosine phosphorylase (xapA).
- deoxyribose-1-phosphate is converted into deoxyribose-5-phosphate through a reaction catalyzed by deoxyribose phosphate mutase (deoB),
- which is further degraded to D-glyceraldehyde-3-phosphate and acetaldehyde through a reaction catalyzed by deoxyribose-5-phosphate aldolase (deoC).

**[0007]** It has been shown that the deo enzymes also catalyze in vitro the reverse anabolic reactions: Deoxyribose-5-phosphate is obtained in vitro in the presence of purified *Escherichia coli* or *Lactobacillus plantarum* deoxyribose aldolase starting from acetaldehyde and D-glyceraldehyde-3-phosphate (Rosen et al., J. Biol. Chem., 240, (1964), 1517-1524; Pricer, J. Biol. Chem., 235, (1960), 1292-1298). Deoxyribose can also be obtained with acetaldehyde and glyceraldehyde as enzyme substrates, but only with a very low yield (Barbas, J. Am. Chem. Soc. 112 (1990), 2013-2014).

**[0008]** The patent application WO 01/14566 describes the enzymatic synthesis of deoxynucleosides starting from deoxyribose-1-phosphate through the combined activities of three enzymes of the deo operon, i.e. deoxyribose aldolase, deoxyribomutase and phosphorylase (thymidine or purine nucleoside phosphorylase) in a one-pot reaction, using as starting substrates glyceraldehyde-3-phosphate, acetaldehyde and a nucleobase. D-glyceraldehyde-3-phosphate can be obtained from fructose-1,6-bisphosphate by an enzymatic process.

**[0009]** The patent application EP 1179598 describes the use of phosphorylase to catalyze the enzymatic production of deoxynucleosides starting from deoxyribose-1-phosphate and nucleobase. The yield of deoxynucleoside synthesis is improved by precipitation of phosphate.

**[0010]** However, methods using enzymes of the deo operon working in the reverse direction compared to their biological function show low yields, which indicates serious drawbacks for their use.

**[0011]** In view of the above-described ineffectiveness of the currently applied processes for producing deoxynucleosides and deoxynucleoside precursors, it is an object of the present invention to provide means and methods for the biosynthetic production of deoxynucleosides and deoxynucleoside precursors starting from cheap and commercially available compounds without being dependent on unreliable natural sources.

**[0012]** In particular, there is a need for alternative methods for the production of deoxynucleosides and deoxynucleoside precursors which allow efficient and economical synthesis of deoxyribonucleosides, by means of which the drawbacks of prior art processes are eliminated.

**[0013]** The present invention relates to a method for producing 2'-deoxynucleosides and precursors thereof starting from 2-dehydro-3-deoxy-D-gluconic acid (KDG) or its salts and comprising a decarboxylation step.

**[0014]** In particular, this method is useful for producing 2-deoxy-D-ribose (DRI) as well as synthetically versatile

enamine derivatives of DRI as 2'-deoxynucleoside precursors.

**[0015]** The decarboxylation step takes place by reacting either KDG or its salts directly, or a derivative of KDG, usually to cleave the C1-C2 bond of the KDG.

**[0016]** In one embodiment of the invention, KDG or one of its salts undergoes (oxidative) decarboxylation leading to 2-deoxy-D-ribonic acid (DRN) or its salts, itself being further converted into 2-deoxy-D-ribose (DRI) or 2-deoxy-D-ribitol (DRL).

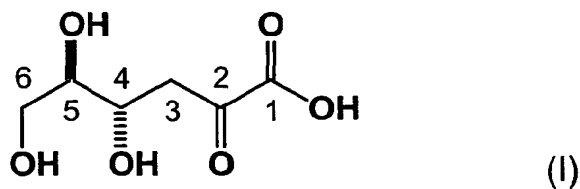
**[0017]** In another embodiment of the invention, decarboxylation takes place by reacting KDG or its salts with an amine, leading to an enamine derivative. This high energy enamine derivative can be further converted into DRI by hydrolysis.

**[0018]** In another embodiment of the invention, (oxidative) decarboxylation is carried out on 3-deoxy-D-gluconic acid (DGN) or its salts and/or 3-deoxy-D-mannonic acid (DMN) or its salts as derivatives of KDG, leading to DRI. Production of a mixture of DGN and DMN takes place by reduction of KDG. The decarboxylation is preferably carried out via reaction with hydrogen peroxide.

**[0019]** In another embodiment of the invention, (oxidative) decarboxylation is carried out on 3-deoxy-D-glucosaminic acid (DGM) or its salts and/or 3-deoxy-D-mannosaminic acid (DMM) or its salts, leading to DRI. Production of a mixture of DGM and DMM takes place from KDG by reductive amination.

**[0020]** Another aspect of the invention is a convenient and cost-effective method for preparing KDG or its salts to be used in the above methods. This method starts either from D-gluconate or from D-glucosamine through the use of recombinant enzymes. The invention provides a novel nucleotide sequence encoding a polypeptide having D-gluconate dehydratase activity and a nucleotide sequence encoding a polypeptide having D-glucosamine deaminase activity.

**[0021]** The starting material used for the method of the present invention is KDG, represented by formula (I) below or one of its salts, or a protected derivative thereof wherein one or more of the hydroxyl groups at positions 4, 5 and/or 6 are protected by a protection group known in the art.



**[0022]** The term "2'-deoxynucleoside" as used herein relates to 2'-deoxyribonucleosides which are N-glycosides, and wherein the basic N-atom of the nucleobase or nucleobase analog is bound to the anomeric carbon atom of 2-deoxy-D-ribose, or one of its derivatives. Examples of a suitable nucleobase are adenine, cytosine, guanine, thymine, uracil, 2,6-diaminopurine, and hypoxanthine. Examples of nucleobase analogs are 5-azacytosine, 2-chloro-adenine, 5-iodo-cytosine, 8-azaguanine, 5-iodo-uracil, 5-bromo-uracil, 5-fluoro-uracil, 5-ethyl-uracil and 5-trifluoromethyl-uracil.

**[0023]** The term "2'-deoxynucleoside precursors" as used herein, relates to compounds which can be easily converted into 2'-deoxynucleosides by applying methods known in the prior art. Preferred 2'-deoxynucleoside precursors are 2-deoxy-D-ribose (DRI) or carbohydrate compounds which can be converted into the 2-deoxy-D-ribosyl moiety of 2'-deoxynucleosides, for instance, those established in the prior art 1-phospho-2-deoxy-D-ribose, 5-phospho-2-deoxy-D-ribose and those established by the present invention 2-deoxy-D-ribitol, 2-deoxy-D-ribonic acid, 2-deoxy-D-ribono-1,4-lactone, 1-N-morpholino-3,4,5-trihydroxy-pentene-1, and their derivatives.

**[0024]** The method of the invention encompasses methods wherein the decarboxylation step is directly carried out on KDG or its salts or on compounds derived from KDG. Preferred KDG derivatives are 3-deoxy-D-gluconic acid, 3-deoxy-D-mannonic acid, 3-deoxy-D-glucosaminic acid and 3-deoxy-D-mannosaminic acid and their respective salts.

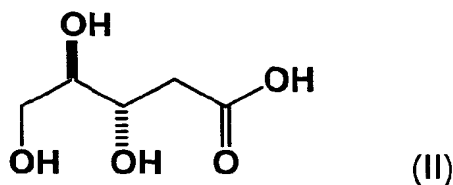
**[0025]** Furthermore, KDG and its salts or protected forms of these wherein one or more of the hydroxyl groups at the positions 4,5 and/or 6 are replaced by protecting groups known for that purpose in the art are also suitable starting materials for the decarboxylation reaction of the present invention. Unless noted otherwise, any reference to KDG in the following specification embraces protected forms of KDG, just as reference to KDG derivatives is intended to embrace protected forms of these derivatives. Similarly, any reference to the products obtained in the methods of the invention is intended to encompass protected forms of these products. Preferred protection groups for the purpose of the invention are those which replace the respective hydroxyl groups by acetate ester, benzoate ester, allyl ether,

benzyl ether, trityl ether, ter-butyldimethylsilyl (TBDMS) ether, isopropylidene or a benzylidene acetal.

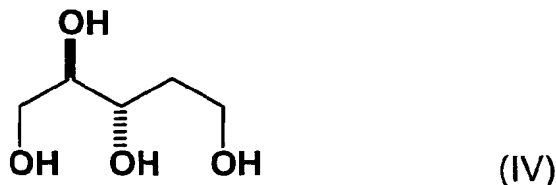
**[0026]** It should be understood that, depending on suitable reaction conditions for the embodiments of the invention, the carboxylic groups contained in the organic acids used as reactants or obtained as products can be in a protonated form or in their salt form, or may be present in equilibrium. Exemplary salts of these acids are those which have metal or ammonium ions as counterions, particularly alkali metal ions such as sodium and/or potassium.

**[0027]** Most of the carbohydrate compounds and their derivatives described in the present invention exist under several cyclic form but for simplicity reasons have been represented by open chain formulas. It is understood that the present invention encompasses all these isomeric or tautomeric forms.

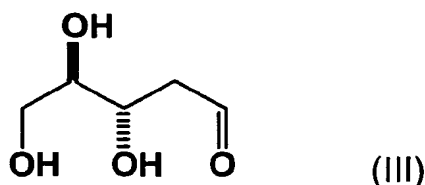
**[0028]** In a first embodiment of the invention, KDG or its salts is reacted with hydrogen peroxide and undergoes (oxidative) decarboxylation to 2-deoxy-D-ribonic acid (DRN), a compound of formula (II) or its salts.



**[0029]** The product may be further converted into or 2-deoxy-D-ribitol (DRL), represented by formula (IV)



or 2-deoxy-D-ribose (DRI), represented by formula (III)



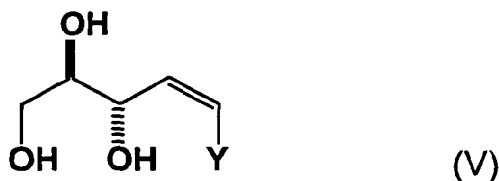
DRN, DRL and particularly DRI are among preferred 2'-deoxynucleoside precursors for the purpose of the present invention. Conversion of DRN to DRI may proceed directly or via DRL as an intermediate.

**[0030]** Preferably, the preparation of DRN is carried out by oxidative decarboxylation of sodium or potassium 2-dehydro-3-deoxy-D-gluconate in aqueous solution with hydrogen peroxide at room temperature as described in example 5. A general method for the preparation of aldonic acids by oxidative decarboxylation of 2-ketoaldonic acids is described in patent EP 1 038 860 A1.

**[0031]** Preferably, the preparation of DRL is carried out by hydrogenation of 2-deoxy-D-ribonolactone in aqueous solution with Rhodium catalyst on carbon at a temperature of 130°C under a pressure of 80 bars as described in example 6. 2-Deoxy-D-ribonolactone can be easily prepared by converting a 2-deoxy-D-ribonate (DRN salt) into 2-deoxy-D-ribonic acid, which is in equilibrium with its lactonic form in aqueous solutions (Han, Tetrahedron. 1993. 49, 349-362; Han, Tetrahedron Asymmetry. 1994. 5, 2535-62).

**[0032]** Preferably the preparation of 2-deoxy-D-ribose (DRI) is carried out by oxidization of 2-deoxy-D-ribitol (DLR), e.g. with chromium oxide in pyridine.

**[0033]** In another embodiment of the invention, decarboxylation takes place by reacting (KDG) or its salts with an amino group-containing reagent Y-H leading to a compound of formula (V).



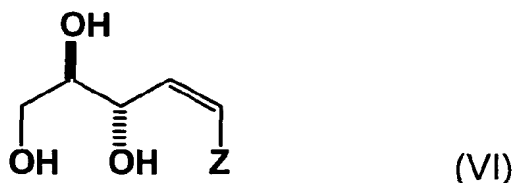
or its respective trans isomer or a protected form thereof, as a 2'-deoxynucleoside precursor. Y-H represents an amine with the hydrogen atom H bound to the nitrogen of the amino group.

**[0034]** In a preferred embodiment of the invention, the amino group-containing reagent represented by Y-H is a linear or cyclic secondary amine; a primary amine that possess a  $\beta$ -carbonyl group, preferably 3-amino-2-indolinone which was found to be effective for the decarboxylation of  $\alpha$ -keto acids (Hanson, J. Chem. Education, 1987, 591-595). In each of these cases, -Y in formula (V) represents the respective nitrogen containing residue derived from these amino-group containing reagent.

**[0035]** Preferably, the compound of formula (V) represents an enamine produced via reaction of a linear or cyclic secondary amine as Y-H.

**[0036]** Preferred cyclic secondary amines are morpholine, pyrrolidine, piperidine, or N-methyl piperazine; preferred non-cyclic amines are those of the formula  $R_1-NH-R_2$ , wherein  $R_1$  and  $R_2$  independently represent a linear or branched alkyl group of 1-8, preferably 1 to 4 carbon atoms. Particularly preferred as a non-cyclic amine is diethylamine. Particularly preferred as a cyclic amine is morpholine.

**[0037]** The compound of formula (V) or its trans isomer or a protected form thereof can be further reacted with Z-H, wherein H represents a hydrogen atom and Z represents a leaving group, to produce a compound of formula (VI)

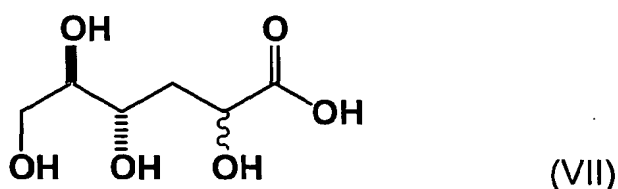


or its respective trans isomer or a protected form thereof, as a 2'-deoxynucleoside precursor. Z-H is preferably water, in which case the compound of formula (VI) is DRI or a protected form thereof (keto-enol-tautomerism).

**[0038]** Preferably, the preparation of the compound of formula (V) is carried out by reacting KDG in benzene with the amine, e.g. morpholine under reflux using the method described in example 7, leading to 1-N-morpholino-3,4,5-tri-hydroxy-pentene-1. Acid catalysed hydrolysis with water yields 2-deoxy-D-ribose (DRI)

**[0039]** A general route to aldehydes via enamines from  $\alpha$ -oxocarboxylic acids carrying  $\beta$ -hydrogens is described by Stamos (Tetrahedron Lett. 23 (1982), 459-462). Other methods for the preparation and hydrolysis of enamines have been described elsewhere (Stork, J. Am. Chem. Soc. 85 (1963), 207-222; Stamhuis, J. Org. Chem. 30 (1965), 2156-2160).

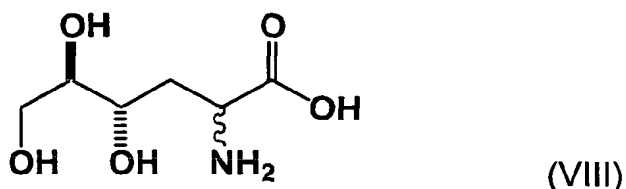
**[0040]** In another embodiment of the invention, KDG or its salt is converted to 3-deoxy-D-gluconic acid (DGN) and/or 3-deoxy-D-mannonic acid (DMN) represented by formula (VII) or the salts of these compounds



**[0041]** The products resulting from this reaction undergo (oxidative) decarboxylation, preferably using hydrogen peroxide, to yield DRI. Production of a mixture of DGN and DMN or their salts takes place from KDG or its salts by reduction.

**[0042]** Preferably the preparation of 2-deoxy-D-ribose (DRI) is carried out by non-stereoselective reduction of 2-dehydro-3-deoxy-D-gluconic acid in water with sodium borohydride at room temperature using the method described for 2-keto-3-deoxyheptonic acid by Weissbach (J. Biol. Chem. 234 (1959), 705-709), followed by oxidative decarboxylation of 3-deoxy-D-gluconate and 3-deoxy-D-mannonate with hydrogen peroxide as described e.g. in US patent 3,312,683; Richards J. Chem. Soc. (1954), 3638-3640; Sowden J. Am. Chem. Soc. 76 (1954), 3541-3542.

**[0043]** In another embodiment of the invention, KDG or its salt is converted to 3-deoxy-D-glucosamine (DGM) or 3-deoxy-D-mannosamine (DMM) represented by formula (VIII) or the salts of these compounds



The products resulting from this reaction undergo (oxidative) decarboxylation, preferably using ninhydrin, to yield DRI. Production of a mixture of DGM and DMM or their salts takes place from KDG or its salts by reductive amination.

**[0044]** Preferably the preparation of 2-deoxy-D-ribose is carried out by non-stereoselective reductive amination of sodium or potassium 2-dehydro-3-deoxy-D-gluconate in aqueous solution with ammonia and sodium cyanoborohydride at room temperature, followed by oxidative decarboxylation of 3-deoxy-D-2-glucosamine and 3-deoxy-D-2-mannosamine with ninhydrin using the method described for the synthesis of 2-deoxy-D-allose by Shelton (J. Am. Chem. Soc. 118 (1996), 2117-2125; and Borch, J. Am. Chem. Soc. 93 (1971), 2897; Durrwachter, J. Am. Chem. Soc. 108 (1986), 7812 referenced therein).

**[0045]** Another aspect of the invention is a convenient and cost-effective method for preparing KDG either from D-gluconate (GCN) or from D-glucosamine through the use of recombinant enzymes.

**[0046]** In a preferred embodiment of the method of the invention, the compound of formula (I) is produced in a preliminary step from a D-gluconate salt by the use of a D-gluconate dehydratase activity. Preferred salts are potassium or sodium D-gluconate. Preferably the D-gluconate dehydratase is encoded by a polynucleotide comprising the nucleotide sequence selected from the group consisting of:

- (a) nucleotide sequences encoding a polypeptide comprising the amino acid sequence of SEQ ID N°2;
- (b) nucleotide sequences comprising the coding sequence of SEQ ID N°1;
- (c) nucleotide sequences encoding a fragment encoded by a nucleotide sequence of (a) or (b);
- (d) nucleotide sequences hybridising with a nucleotide sequence of any one of (a) to (c); and
- (e) nucleotide sequences which deviate from the nucleoside sequence of (d) as a result of degeneracy of the genetic code.

**[0047]** The enzymatic synthesis of KDG or its salts using D-gluconate dehydratase proceeds according to the following reaction: D-gluconate is converted into KDG by the elimination of one water molecule. The activity of a D-gluconate dehydratase has been characterized in different bacteria species e.g. in *Alcaligenes* (Kerstens, Methods in Enzymology 42 (1975), 301-304); *Clostridium pasteurianum*, (Gottschalk, Methods in Enzymology 90 (1982), 283-287); *Thermoplasma acidophilum* (Budgen, FEBS Letters 196 (1986), 207-210) and *Sulfolobus solfataricus* (Nicolaus, Bio-

technology Letters 8(7) (1986), 497-500). The preferred D-gluconate dehydratase was identified by screening several collection strains for D-gluconate dehydratase activity. The gene encoding a D-gluconate dehydratase, which was designated gcnD was selected from a genomic library of *Agrobacterium tumefaciens* strain C58, and further inserted in a multi copy vector optimised for expression. It was shown that a crude extract from *E. coli* cells over-expressing the gcnD gene catalysed the total conversion of D-gluconate into KDG (see Example 2).

**[0048]** In a further preferred embodiment of the method of the invention, the compound of formula (I) is produced in a preliminary step from D-glucosamine by the use of a D-glucosamine deaminase activity. Preferably the D-glucosamine deaminase is encoded by a polynucleotide comprising the nucleotide sequence selected from the group consisting of:

- (f) nucleotide sequences encoding a polypeptide comprising the amino acid sequence of SEQ ID N°4;
- (g) nucleotide sequences comprising the coding sequence of SEQ ID N°3;
- (h) nucleotide sequences encoding a fragment encoded by a nucleotide sequence of (a) or (b);
- (i) nucleotide sequences hybridising with a nucleotide sequence of any one of (a) to (c); and
- (j) nucleotide sequences which deviate from the nucleoside sequence of (d) as a result of degeneracy of the genetic code.

**[0049]** The enzymatic synthesis of KDG or its salts using D-glucosamine deaminase proceeds according to the following reaction: D-glucosamine is converted into KDG by the elimination of one molecule water and one molecule of ammonia. The activity of a D-glucosamine deaminase has been characterized in different bacteria species e.g. in *Pseudomonas fluorescens* (Iwamoto, Agric. Biol. Chem. 53 (1989), 2563-2569) *Agrobacterium radiobacter* (Iwamoto, FEBS Letters 104 (1979), 131-134; Iwamoto, J. Biochem. 91 (1982), 283-289), and its requirement for Mn<sup>2+</sup> ion was shown (Iwamoto, Biosci. Biotech. Biochem. 59 (1995), 408-411).

**[0050]** The preferred D-glucosamine deaminase was identified by screening several collection strains for D-glucosamine deaminase activity. The gene encoding a D-glucosamine deaminase, which was designated gmaA was isolated from *Agrobacterium tumefaciens* strain C58 by cloning a gene annotated as a putative D-serine deaminase. The gmaA gene was further inserted in a multi copy vector optimised for expression. It was shown that a crude extract from *E. coli* cells over-expressing the gmaA gene catalysed the conversion of D-glucosamine into KDG (see Example 4).

**[0051]** These and other embodiments are disclosed and encompassed by the description and examples of the present invention. Further literature concerning any one of the methods, uses and compounds to be employed in accordance with the present invention may be retrieved from public libraries, using for example electronic devices.

**[0052]** For example the public database "Medline" may be utilized which is available on the Internet, for example under <http://www.ncbi.nlm.nih.gov/PubMed/medline.html>. Further databases and addresses, such as <http://www.ncbi.nlm.nih.gov/>, <http://www.infobiogen.fr/>, [http://www.fmi.ch/biology/research\\_tools.html](http://www.fmi.ch/biology/research_tools.html), <http://www.tigr.org/>, are known to the person skilled in the art and can also be obtained using, e.g., <http://www.google.de>. An overview of patent information in biotechnology and a survey of relevant sources of patent information useful for retrospective searching and for current awareness is given in Berks, TIBTECH 12 (1994), 352-364.

Furthermore, the term "and/or" when occurring herein includes the meaning of "and", "or" and "all or any other combination of the elements connected by said term".

## EXAMPLES

### Example 1. Cloning of a gene encoding a D-gluconate dehydratase from *Agrobacterium tumefaciens* strain C58 (CIP 104333)

**[0053]** *Agrobacterium tumefaciens* strain C58 (CIP 104333) was obtained from Institut Pasteur Collection (CIP, Paris, France). Chromosomal DNA was extracted and a D-gluconate dehydratase gene was amplified by PCR according to standard protocols using the following primers:

5'-CCCTTAATTAATGACGACATCTGATAATCTTC-3', depicted in SEQ ID N° 5;

5'-TTTGCGGCCGCTTAGTGGTTATCGCGCGGC-3', depicted in SEQ ID N° 6;

5'-CCCGGTACCATGACGACATCTGATAATCTTC-3', depicted in SEQ ID N° 7;

A first DNA fragment amplified using the two primers depicted in SEQ ID N° 5 and SEQ ID N° 6, was ligated into a pUC18-derived vector previously digested by *PacI* and *NotI* to yield the plasmid pVDM80. A second DNA fragment amplified using the two primers depicted in SEQ ID N° 6 and SEQ ID N° 7, was ligated into a pET29a vector (Novagen) previously digested by *KpnI* and *NotI* to yield the plasmid pVDM82. The nucleotide sequence of the cloned gene is depicted in SEQ ID N° 1 and the sequence of the polypeptide encoded by this gene is depicted in SEQ ID N° 2.

**Example 2. Expression of a D-gluconate dehydratase activity in Escherichia coli and preparation of 2-dehydro-3-deoxy-D-gluconate from D-gluconate.**

[0054] Competent cells of *E. coli* BL21 were transformed with the pVDM82 plasmid constructed as described in example 1 yielding strain +1289. Strain + 1289 was cultivated at 30°C in Luria-Bertani (LB) medium (Difco) containing 30 mg/l kanamycin until OD(600 nm) reached a value of 0.6. Then isopropyl- $\beta$ -D-thiogalactopyranoside (IPTG) was added to a 0.5 mM final concentration. After a further cultivation period of 2 hours and 30 minutes, cells were collected by centrifugation and washed once with 20 mM sodium phosphate buffer pH 7.2. A cell extract was prepared by suspending about 5 g of cells in 10 ml of Tris-HCl 50 mM pH 8.5 buffer containing 10000 units lysozyme (Ready-Lyse, Epicentre, Madison, Wisconsin) and 1 mM EDTA, and incubating the suspension at 30°C for 15 minutes. Then 10000 kUnits deoxyribonuclease I (DNase I, Sigma) as well as 5 mM MgCl<sub>2</sub> were added to the preparation which was incubated at 30°C for an additional period of 15 minutes. The cell extract thus obtained was kept frozen at -20°C before use. 1.5 ml of the cell extract was mixed with 2M sodium or potassium D-gluconate in a total volume of 10 ml. This preparation was incubated at 37°C after the pH has been adjusted to 8.5. The progression of 2-dehydro-3-deoxy-D-gluconate (KDG) synthesis was followed by analysing aliquots taken after increasing periods of incubation. Several dilution parts of these aliquots were deposited on silica plates and chromatographed in the following solvent system: isopropanol / water (90/10). A yellow spot of KDG (R<sub>f</sub> ~0.40) was detected after revelation with p-anisaldehyde. KDG was also quantitated using a spectrophotometric assay based on the reaction with semicarbazide hydrochloride as described by Mac Gee (J. Biol. Chem. 1954. 210, 617-626). Typically, after a 30h period of incubation and using the spectrophotometric assay, KDG concentration ranged from 1.5 to 2 M.

The sodium or potassium 2-dehydro-3-deoxy-D-gluconate solution thus obtained could be used as such for further synthetic steps. 2-Dehydro-3-deoxy-D-gluconic acid could also be prepared from such a solution applying published protocols (Bender, Anal. Biochem. 1974. 61, 275-279). A crude preparation of a mixture of 2-dehydro-3-deoxy-D-gluconic acid and KCl could also be obtained by adding one equivalent of HCl to a potassium 2-dehydro-3-deoxy-D-gluconate solution which was then evaporated.

**Example 3. Cloning of a gene encoding a D-glucosamine deaminase from Agrobacterium tumefaciens strain C58 (CIP 104333)**

[0055] *Agrobacterium tumefaciens* strain C58 (CIP 104333) was obtained from Institut Pasteur Collection (CIP, Paris, France). Chromosomal DNA was extracted and a D-glucosamine deaminase gene was amplified by PCR according to standard protocols using the following primers:

5'-CCCTTAATTAATGCAGTCTTCTTCAGCTCTTC-3', depicted in SEQ ID N° 8;

5'-TTTGCGGCCCGCCTAGTGAAAGAAGGTTGTGTAGAT-3', depicted in SEQ ID N° 9;

5'-AAATCATGACTATGCAGTCTTCTTCAGCTCTTCG-3', depicted in SEQ ID N° 10;

5'-TATAGATCTCTAGTGAAAGAAGGTTGTGTAGAT-3', depicted in SEQ ID N° 11;

A first DNA fragment amplified using the two primers depicted in SEQ ID N° 8 and SEQ ID N° 9, was ligated into a pUC18-derived vector previously digested by PaeI and NotI to yield the plasmid pKDGb1. A second DNA fragment amplified using the two primers depicted in SEQ ID N° 10 and SEQ ID N° 11, was ligated into a pQE60 vector (Qiagen) previously digested by BspH1 and BglII to yield the plasmid pEP18. The nucleotide sequence of the cloned gene is depicted in SEQ ID N° 3 and the sequence of the polypeptide encoded by this gene is depicted in SEQ ID N° 4.

**Example 4. Expression of a D-glucosamine deaminase activity in Escherichia coli and preparation of 2-dehydro-3-deoxy-D-gluconic acid from D-glucosamine**

[0056] Competent cells of *E. coli* MG1655 were transformed with the pEP18 plasmid constructed as described in example 1 and pREP4 (Qiagen) yielding strain +1068. Strain + 1068 was cultivated at 37°C in LB medium containing 30 mg/l kanamycin and 100 mg/l ampicillin until OD(600 nm) reached a value of 0.6. Then IPTG was added to a 0.5 mM final concentration. After a further cultivation period of 2 hours and 30 minutes, cells were collected by centrifugation and washed once with 20 mM sodium phosphate buffer pH 7.2. A cell extract was prepared using the protocol described in example 2.

[0057] 2 ml of the cell extract was mixed with 100 mM sodium or potassium D-glucosamine and 0.1 mM pyridoxal phosphate in a total volume of 5 ml. This preparation was incubated at 37°C after the pH has been adjusted to 7.5.

The progression of 2-dehydro-3-deoxy-D-gluconate (KDG) synthesis was followed using the protocols described in example 2. Typically, after a 30h period of incubation and using the spectrophotometric assay described in example 2, KDG concentration ranged from 50 to 100 mM.



**Example 5. Preparation of 2-deoxy-D-ribonate from 2-dehydro-3-deoxy-D-gluconate**

**[0058]** 0.5 ml of a 31% hydrogen peroxyde solution were added to 5 ml of a 1M potassium 2-dehydro-3-deoxy-D-gluconate (KDG) solution at 25°C. The progression of KDG decarboxylation was followed both by the observation of bubbles resulting from the release of carbon dioxide and by the disappearance of KDG using the thin layer chromatography protocol described in example 2. Typically, after a 3h period of reaction the concentration of residual KDG was less than 10 mM.

**Example 6. Preparation of 2-deoxy-D-ribitol from 2-deoxy-D-ribonolactone**

**[0059]** 0.2 g of Rhodium ( 5 % on carbon) catalyst was added to an aqueous solution of 1 g 2-deoxy-D-ribonolactone prepared following a method described by Deriaz (J. Chem. Soc. (1949), 1879-1883) for the synthesis of 2-deoxy-L-ribonolactone. Hydrogenation of 2-deoxy-D-ribonolactone was performed at 130°C under a pressure of 80 bars. The solution obtained after filtration of the reaction mixture was evaporated. The residue was dissolved in ethyl acetate and further purified by chromatography on a silica column. The solvent was removed in vacuo leading to a yellow oil (yield 85%). The compound thus obtained was identical with 2-deoxy-D-ribitol obtained by reduction of 2-deoxy-D-ribose as described by Rabow (J. Am. Chem. Soc. 122 (1999), 3196-3203).

**Example 7. Preparation of 1-N-morpholino-3,4,5-trihydroxypentene-1 from 2-dehydro-3-deoxy-D-gluconate.**

**[0060]** 2 g of 2-dehydro-3-deoxy-D-gluconic acid were suspended in 150 ml benzene. 1.1 ml morpholine and 100 mg p-toluenesulfonic acid were added to the suspension and the reaction mixture was refluxed for 3 hours. Water formed by this reaction was removed by distillation. Benzene was decanted. Solid compounds attached to the vessel were collected, washed with acetone and dried. The main compound present in this preparation (yield 40%) was further purified by column chromatography on a silica column using a gradient of methanol in chloroform. Fractions containing 1-N-morpholino-3,4,5-trihydroxypentene-1 were pooled and solvent was removed in vacuo. <sup>1</sup>H-NMR (D<sub>2</sub>O): δ = 3.15 ppm (4H, t, morpholine), 3.8 ppm (4H, t, morpholine), 3.4 to 4 ppm, (4H, m, 5a-H, 5b-H, 4-H, 3-H), 6.3 and 6.8 ppm (2H, 2d, 1-H and 2-H, J = 4 Hz).

## SEQUENCE LISTING

5 <110> Evologic S.A., MARLIERE Philippe

<120> PRODUCTION OF 2'-DEOXYNUCLEOSIDES AND 2'-DEOXYNUCLEOSIDE PRECURSORS  
FROM 2-DEHYDRO-3-DEOXY-D-GLUCONATE

10 <130> G 3111 EP

15 <160> 11

<170> PatentIn version 3.1

20 <210> 1

<211> 1812

25 <212> DNA

<213> Agrobacterium tumefaciens

<220>

30 <221> CDS

<222> (1)..(1809)

35 <223>

<400> 1

atg acg aca tct gat aat ctt cct gca act cag ggc aag ctc cgt tcg	48
Met Thr Thr Ser Asp Asn Leu Pro Ala Thr Gln Gly Lys Leu Arg Ser	
1 5 10 15	
cgc gcc tgg ttc gac aac cca gcc aat gcg gac atg acc gcg ctt tat	96
Arg Ala Trp Phe Asp Asn Pro Ala Asn Ala Asp Met Thr Ala Leu Tyr	
20 25 30	
ctc gag cgt tac atg aac ttc ggt ctc agc cag gcc gag ctt cag tcc	144
Leu Glu Arg Tyr Met Asn Phe Gly Leu Ser Gln Ala Glu Leu Gln Ser	
35 40 45	
gac cgc ccg att atc ggt att gcg cag acc ggt tcc gac ctt tcg ccc	192
Asp Arg Pro Ile Ile Gly Ile Ala Gln Thr Gly Ser Asp Leu Ser Pro	
50 55 60	
tgc aac cgt cat cat ctg gaa ctc gcc aac cgt ctg cgt gaa ggc att	240
Cys Asn Arg His His Leu Glu Leu Ala Asn Arg Leu Arg Glu Gly Ile	
65 70 75 80	
cgt gaa gcc ggc ggc atc gcc atc gaa ttc ccg gtg cat ccg atc cag	288
Arg Glu Ala Gly Gly Ile Ala Ile Glu Phe Pro Val His Pro Ile Gln	
85 90 95	

55

EP 1 491 549 A1

		gaa acc ggt aag cgt ccg aca gcg ggc ctt gat cgc aac ctg gct tac	336
		Glu Thr Gly Lys Arg Pro Thr Ala Gly Leu Asp Arg Asn Leu Ala Tyr	
		100 105 110	
5		ctc ggc ctc gtg gaa gtg ctt tat ggc tat ccg ctc gac ggc gtt gtt	384
		Leu Gly Leu Val Glu Val Leu Tyr Gly Tyr Pro Leu Asp Gly Val Val	
		115 120 125	
		ctg acc atc ggc tgc gac aag acc acg cct gcc tgt ctt atg gcg gcg	432
		Leu Thr Ile Gly Cys Asp Lys Thr Thr Pro Ala Cys Leu Met Ala Ala	
		130 135 140	
10		gcc acc gtc aac att ccg gcc atc gcc ctt tcc gtc ggt ccc atg ctg	480
		Ala Thr Val Asn Ile Pro Ala Ile Ala Leu Ser Val Gly Pro Met Leu	
		145 150 155	
		aac ggc tgg ttc cgc ggt gag cgc acc ggt tcc gcc acc atc gtc tgg	528
		Asn Gly Trp Phe Arg Gly Glu Arg Thr Gly Ser Gly Thr Ile Val Trp	
		165 170 175	
		aag gcc cgc gaa ctg ctg gcg aag ggc gag atc gat tac cag ggc ttc	576
		Lys Ala Arg Glu Leu Leu Ala Lys Gly Glu Ile Asp Tyr Gln Gly Phe	
		180 185 190	
20		gtc aag ctc gtt gcc tgc tct gcc ccg tcc acc ggc tat tgc aac acc	624
		Val Lys Leu Val Ala Ser Ser Ala Pro Ser Thr Gly Tyr Cys Asn Thr	
		195 200 205	
		atg ggc acg gca aca acc atg aac tgc ctc gcc gaa gcg ctc ggc atg	672
		Met Gly Thr Ala Thr Thr Met Asn Ser Leu Ala Glu Ala Leu Gly Met	
		210 215 220	
25		cag ctt ccc ggc tcc gcc gcc att ccg gcg cct tac cgt gac cgt cag	720
		Gln Leu Pro Gly Ser Ala Ala Ile Pro Ala Tyr Arg Asp Arg Gln	
		225 230 235 240	
		gaa gtc tct tac ctc acc ggc ctg cgc atc gtc gac atg gtc agg gaa	768
		Glu Val Ser Tyr Leu Thr Gly Leu Arg Ile Val Asp Met Val Arg Glu	
		245 250 255	
30		gac ctg aaa cca tca gac atc atg acc aag gat gcc ttc atc aac gcc	816
		Asp Leu Lys Pro Ser Asp Ile Met Thr Lys Asp Ala Phe Ile Asn Ala	
		260 265 270	
35		atc cgc gtt aat tgc gcg atc ggc ggt tcc acc aac gcg ccg atc cat	864
		Ile Arg Val Asn Ser Ala Ile Gly Ser Thr Asn Ala Pro Ile His	
		275 280 285	
		cta aac ggc ctt gcc cgc cat gtc ggc gtc gag ctg acg gtg gat gac	912
		Leu Asn Gly Leu Ala Arg His Val Gly Val Glu Leu Thr Val Asp Asp	
		290 295 300	
40		tgg cag acc tat ggc gaa gac gtg ccg ctg ctc gtc aac ctg cag ccg	960
		Trp Gln Thr Tyr Gly Glu Asp Val Pro Leu Leu Val Asn Leu Gln Pro	
		305 310 315 320	
		gca ggc gaa tat ctc ggc gag gac tat tac cat gcc ggc ggc gtt ccc	1008
		Ala Gly Glu Tyr Leu Gly Glu Asp Tyr Thr His Ala Gly Gly Val Pro	
		325 330 335	
45		gct gtc gtc aat cag ctg atg acc caa ggg ctg atc atg gaa gac gcc	1056
		Ala Val Val Asn Gln Leu Met Thr Gln Gly Leu Ile Met Glu Asp Ala	
		340 345 350	
50		atg acc gtc aac ggc aag acc atc ggc gac aat tgc cgt ggc gcg atc	1104
		Met Thr Val Asn Gly Lys Thr Ile Gly Asp Asn Cys Arg Gly Ala Ile	
		355 360 365	
		atc gaa gac gag aag gtc atc cgc ccc tat gag cag ccg ctc aag gag	1152
		Ile Glu Asp Glu Lys Val Ile Arg Pro Tyr Glu Gln Pro Leu Lys Glu	
55			

# EP 1 491 549 A1

	370	375	380	
5	cgt gcc ggc ttc cgc gtt ctg cgc ggc aat ctg ttc tcc tcg gcc atc Arg Ala Gly Phe Arg Val Leu Arg Gly Asn Leu Phe Ser Ser Ala Ile 385 390 395 400			1200
	atg aag aca agc gtg att tcg gaa gaa ttc cgc ggt cgt tac ctc tcc Met Lys Thr Ser Val Ile Ser Glu Glu Phe Arg Gly Arg Tyr Leu Ser 405 410 415			1248
10	aac cct gat gat ccg gaa gcc ttc gaa ggc cgc gcc gtg gtg ttc gat Asn Pro Asp Asp Pro Glu Ala Phe Glu Gly Arg Ala Val Val Phe Asp 420 425 430			1296
	ggg ccg gag gat tac cat cat cgc atc gac gat ccg tcg ctt ggc atc Gly Pro Glu Asp Tyr His His Arg Ile Asp Asp Pro Ser Leu Gly Ile 435 440 445			1344
15	gac gcc aac acc gtc ctg ttc atg cgc ggc gcc ggt ccg atc ggt tac Asp Ala Asn Thr Val Leu Phe Met Arg Gly Ala Gly Pro Ile Gly Tyr 450 455 460			1392
20	ccg ggc gca gcg gaa gtg gtg aac atg cgc gcg ccg gat tac ctt ctg Pro Gly Ala Ala Glu Val Val Asn Met Arg Ala Pro Asp Tyr Leu Leu 465 470 475 480			1440
	aag caa ggc gtc agt tcg ctg ccc tgc atc ggc gat ggc cgc cag tcc Lys Gln Gly Val Ser Ser Leu Pro Cys Ile Gly Asp Gly Arg Gln Ser 485 490 495			1488
25	ggc acg tcg ggc agc cca tcc atc ctc aat gcc tcg ccg gaa gcg gcg Gly Thr Ser Gly Ser Pro Ser Ile Leu Asn Ala Ser Pro Glu Ala Ala 500 505 510			1536
	gcc ggc ggc ggt ctg tct att ctg cag acg ggt gac cgc gtc cgc atc Ala Gly Gly Gly Leu Ser Ile Leu Gln Thr Gly Asp Arg Val Arg Ile 515 520 525			1584
30	gat gtg ggc cgc ggc aag gcc gat atc ctg ata tca ggt gaa gag ctc Asp Val Gly Arg Gly Lys Ala Asp Ile Leu Ile Ser Gly Glu Glu Leu 530 535 540			1632
	gcc aag cgt tac gag gcg ctg gca gct cag ggc ggt tat aag ttc ccc Ala Lys Arg Tyr Glu Ala Leu Ala Ala Gln Gly Gly Tyr Lys Phe Pro 545 550 555 560			1680
35	gac cac cag acg ccg tgg cag gaa atc cag cgc ggt atc gtc agc cag Asp His Gln Thr Pro Trp Gln Glu Ile Gln Arg Gly Ile Val Ser Gln 565 570 575			1728
40	atg gaa acc ggc gcg gtt ctg gaa ccg gcc gta aag tat cag cgc atc Met Glu Thr Gly Ala Val Leu Glu Pro Ala Val Lys Tyr Gln Arg Ile 580 585 590			1776
	gcc cag acc aag ggc ctg ccg cgc gat aac cac tga Ala Gln Thr Lys Gly Leu Pro Arg Asp Asn His 595 600			1812
45	<210> 2			
	<211> 603			
	<212> PRT			
50	<213> Agrobacterium tumefaciens			
	<400> 2			
55				

EP 1 491 549 A1

5 Met Thr Thr Ser Asp Asn Leu Pro Ala Thr Gln Gly Lys Leu Arg Ser  
 1 5 10 15  
 Arg Ala Trp Phe Asp Asn Pro Ala Asn Ala Asp Met Thr Ala Leu Tyr  
 20 25 30  
 10 Leu Glu Arg Tyr Met Asn Phe Gly Leu Ser Gln Ala Glu Leu Gln Ser  
 35 40 45  
 Asp Arg Pro Ile Ile Gly Ile Ala Gln Thr Gly Ser Asp Leu Ser Pro  
 50 55 60  
 15 Cys Asn Arg His His Leu Glu Leu Ala Asn Arg Leu Arg Glu Gly Ile  
 65 70 75 80  
 Arg Glu Ala Gly Gly Ile Ala Ile Glu Phe Pro Val His Pro Ile Gln  
 85 90 95  
 20 Glu Thr Gly Lys Arg Pro Thr Ala Gly Leu Asp Arg Asn Leu Ala Tyr  
 100 105 110  
 Leu Gly Leu Val Glu Val Leu Tyr Gly Tyr Pro Leu Asp Gly Val Val  
 115 120 125  
 25 Leu Thr Ile Gly Cys Asp Lys Thr Thr Pro Ala Cys Leu Met Ala Ala  
 130 135 140  
 30 Ala Thr Val Asn Ile Pro Ala Ile Ala Leu Ser Val Gly Pro Met Leu  
 145 150 155 160  
 Asn Gly Trp Phe Arg Gly Glu Arg Thr Gly Ser Gly Thr Ile Val Trp  
 165 170 175  
 35 Lys Ala Arg Glu Leu Leu Ala Lys Gly Glu Ile Asp Tyr Gln Gly Phe  
 180 185 190  
 Val Lys Leu Val Ala Ser Ser Ala Pro Ser Thr Gly Tyr Cys Asn Thr  
 195 200 205  
 40 Met Gly Thr Ala Thr Thr Met Asn Ser Leu Ala Glu Ala Leu Gly Met  
 210 215 220  
 45 Gln Leu Pro Gly Ser Ala Ala Ile Pro Ala Pro Tyr Arg Asp Arg Gln  
 225 230 235 240  
 Glu Val Ser Tyr Leu Thr Gly Leu Arg Ile Val Asp Met Val Arg Glu  
 245 250 255  
 50 Asp Leu Lys Pro Ser Asp Ile Met Thr Lys Asp Ala Phe Ile Asn Ala  
 260 265 270  
 55 Ile Arg Val Asn Ser Ala Ile Gly Gly Ser Thr Asn Ala Pro Ile His

EP 1 491 549 A1

	275	280	285
5	Leu Asn Gly Leu Ala Arg His Val Gly Val Glu Leu Thr Val Asp Asp 290 295 300		
	Trp Gln Thr Tyr Gly Glu Asp Val Pro Leu Leu Val Asn Leu Gln Pro 305 310 315 320		
10	Ala Gly Glu Tyr Leu Gly Glu Asp Tyr Tyr His Ala Gly Gly Val Pro 325 330 335		
	Ala Val Val Asn Gln Leu Met Thr Gln Gly Leu Ile Met Glu Asp Ala 340 345 350		
15	Met Thr Val Asn Gly Lys Thr Ile Gly Asp Asn Cys Arg Gly Ala Ile 355 360 365		
	Ile Glu Asp Glu Lys Val Ile Arg Pro Tyr Glu Gln Pro Leu Lys Glu 370 375 380		
20	Arg Ala Gly Phe Arg Val Leu Arg Gly Asn Leu Phe Ser Ser Ala Ile 385 390 395 400		
	Met Lys Thr Ser Val Ile Ser Glu Glu Phe Arg Gly Arg Tyr Leu Ser 405 410 415		
25	Asn Pro Asp Asp Pro Glu Ala Phe Glu Gly Arg Ala Val Val Phe Asp 420 425 430		
30	Gly Pro Glu Asp Tyr His His Arg Ile Asp Asp Pro Ser Leu Gly Ile 435 440 445		
	Asp Ala Asn Thr Val Leu Phe Met Arg Gly Ala Gly Pro Ile Gly Tyr 450 455 460		
35	Pro Gly Ala Ala Glu Val Val Asn Met Arg Ala Pro Asp Tyr Leu Leu 465 470 475 480		
40	Lys Gln Gly Val Ser Ser Leu Pro Cys Ile Gly Asp Gly Arg Gln Ser 485 490 495		
	Gly Thr Ser Gly Ser Pro Ser Ile Leu Asn Ala Ser Pro Glu Ala Ala 500 505 510		
45	Ala Gly Gly Gly Leu Ser Ile Leu Gln Thr Gly Asp Arg Val Arg Ile 515 520 525		
	Asp Val Gly Arg Gly Lys Ala Asp Ile Leu Ile Ser Gly Glu Glu Leu 530 535 540		
50	Ala Lys Arg Tyr Glu Ala Leu Ala Ala Gln Gly Gly Tyr Lys Phe Pro 545 550 555 560		
55			

# EP 1 491 549 A1

Asp His Gln Thr Pro Trp Gln Glu Ile Gln Arg Gly Ile Val Ser Gln  
 565 570 575  
 5  
 Met Glu Thr Gly Ala Val Leu Glu Pro Ala Val Lys Tyr Gln Arg Ile  
 580 585 590  
 Ala Gln Thr Lys Gly Leu Pro Arg Asp Asn His  
 595 600  
 10  
 <210> 3  
 <211> 1272  
 15  
 <212> DNA  
 <213> Agrobacterium tumefaciens  
 <220>  
 20  
 <221> CDS  
 <222> (1)..(1269)  
 <223>  
 25  
 <400> 3  
 atg cag tct tct tca gct ctt cgg caa tca acc ggc gat cag tcg gaa 48  
 Met Gln Ser Ser 5 Ala Leu Arg Gln Ser Thr Gly Asp Gln Ser Glu  
 1 10 15  
 30  
 tac cat gcc cag tcg aat atg atc ggc tct agc ccg gcg gac ggt ttg 96  
 Tyr His Ala Gln Ser Asn Met Ile Gly Ser Ser Pro Ala Asp Gly Leu  
 20 25 30  
 35  
 ctc gca ttg ccg ctt ctg acc gtc gat ctt gcc gtc tat cgc ggt aat 144  
 Leu Ala Leu Pro Leu Leu Thr Val Asp Leu Ala Val Tyr Arg Gly Asn  
 35 40 45  
 cgg gat cgc ttt ctt gcg ctt gtc tcg gcc cat gga gcg aag gcg gct 192  
 Arg Asp Arg Phe Leu Ala Leu Val Ser Ala His Gly Ala Lys Ala Ala  
 50 55 60  
 40  
 cca cat gcc aag acg ccg atg tgc ccg gag atc gcg atc gat ctg att 240  
 Pro His Ala Lys Thr Pro Met Cys Pro Glu Ile Ala Ile Asp Leu Ile  
 65 70 75 80  
 45  
 gaà gcc ggt gcc tgg ggc gcg acg gtc gcc gat ctc ttc cag gcg gaa 288  
 Glu Ala Gly Ala Trp Gly Ala Thr Val Ala Asp Leu Phe Gln Ala Glu  
 85 90 95  
 50  
 gtc ctg ctc aag gcc ggc gtg tcg aac ata ttg atc gcc aac cag atc 336  
 Val Leu Leu Lys Ala Gly Val Ser Asn Ile Leu Ile Ala Asn Gln Ile  
 100 105 110  
 ggc gga ttg aca tcc gcc aga cgc cta cgc atg ctc gca gat gct ttt 384  
 Gly Gly Leu Thr Ser Ala Arg Arg Leu Arg Met Leu Ala Asp Ala Phe  
 115 120 125  
 55  
 ccg aaa gcc gag att atc tgc tgt gtc gat tct gtt cag gcc tcg gcc 432  
 Pro Lys Ala Glu Ile Ile Cys Cys Val Asp Ser Val Gln Ala Ser Ala  
 130 135 140

EP 1 491 549 A1

	aat ctg gtt cag gcc ttt caa ggg cgt gtg gat gcc cca ttc aag gtc Asn Leu Val Gln Ala Phe Gln Gly Arg Val Asp Ala Pro Phe Lys Val 145 150 155 160	480
5	ttc atc gaa gtc ggt gtc ggc cgc act ggc gcc cgt acg ttg aat gtt Phe Ile Glu Val Gly Val Gly Arg Thr Gly Ala Arg Thr Leu Asn Val 165 170 175	528
10	gca aag gat atc atc gac acc atc tcg aca agt gca gaa atc gta ctg Ala Lys Asp Ile Ile Asp Thr Ile Ser Thr Ser Ala Glu Ile Val Leu 180 185 190	576
	gcc ggt gtg tcg acc tat gaa ggc tcc gtc tcc ggg gaa acg tcg gaa Ala Gly Val Ser Thr Tyr Glu Gly Ser Val Ser Gly Glu Thr Ser Glu 195 200 205	624
15	gca ctc gat gca aac atg gcg gcc ctg ttc gat ctc ctg acc gac agt Ala Leu Asp Ala Asn Met Ala Ala Leu Phe Asp Leu Leu Thr Asp Ser 210 215 220	672
	ctt gca tcg ata cgc gaa aaa gat ccc ggg cgc ccg cta acg gtt tca Leu Ala Ser Ile Arg Glu Lys Asp Pro Gly Arg Pro Leu Thr Val Ser 225 230 235 240	720
20	gcc ggc ggt tcg atc cat ttc gac cgc gtg ctc gcg gcg ctt gtg ccc Ala Gly Gly Ser Ile His Phe Asp Arg Val Leu Ala Ala Leu Val Pro 245 250 255	768
25	gtt tgc gag gcg gat ggc aat gcg acg ttg ttg ctg cgc agc ggc gcc Val Cys Glu Ala Asp Gly Asn Ala Thr Leu Leu Leu Arg Ser Gly Ala 260 265 270	816
	atc ttc ttc tct gat cac ggt gta tat cag cgc ggt ttc cag gca gtc Ile Phe Phe Ser Asp His Gly Val Tyr Gln Arg Gly Phe Gln Ala Val 275 280 285	864
30	gac gcc cgc aac cta ctc gca tcc ggc aag gtt gtc ttc aag gca tcc Asp Ala Arg Asn Leu Leu Ala Ser Gly Lys Val Val Phe Lys Ala Ser 290 295 300	912
	gag gca ttt cag ccc tca atg cga atc tgg gcg gag gtc atc tcc gtt Glu Ala Phe Gln Pro Ser Met Arg Ile Trp Ala Glu Val Ile Ser Val 305 310 315 320	960
35	cct gag ccg ggg ctg gcg atc gtc ggc atg ggc atg ccg gat gta tcg Pro Glu Pro Gly Leu Ala Ile Val Gly Met Gly Met Arg Asp Val Ser 325 330 335	1008
40	ttc gat cag gac ctg ccc gtg gcg ctt cgg ctc cat agg gac gga cat Phe Asp Gln Asp Leu Pro Val Ala Leu Arg Leu His Arg Asp Gly His 340 345 350	1056
	ctg gtc gaa gct gat ctc tct tca tcc gcg aag gtc ggc aag ctc aat Leu Val Glu Ala Asp Leu Ser Ser Ser Ala Lys Val Gly Lys Leu Asn 355 360 365	1104
45	gac cag cat gcc ttc ttg tcc ttc ggg aac ggc agc agt ctg gca atc Asp Gln His Ala Phe Leu Ser Phe Gly Asn Gly Ser Ser Leu Ala Ile 370 375 380	1152
	ggc gat gtc ata gaa ttc ggc atc tcg cat ccc tgc acg tgc ttc gat Gly Asp Val Ile Glu Phe Gly Ile Ser His Pro Cys Thr Cys Phe Asp 385 390 395 400	1200
50	cgc tgg cgc gtc ttt cac gga atc gat gga tca ggc cgg atc cag cgc Arg Trp Arg Val Phe His Gly Ile Asp Gly Ser Gly Arg Ile Gln Arg 405 410 415	1248
55	atc tac aca acc ttc ttt cac tag Ile Tyr Thr Thr Phe Phe His	1272



EP 1 491 549 A1

420

5 <210> 4  
 <211> 423  
 <212> PRT  
 <213> Agrobacterium tumefaciens

10

<400> 4

Met Gln Ser Ser Ser Ala Leu Arg Gln Ser Thr Gly Asp Gln Ser Glu  
 1 5 10 15

Tyr His Ala Gln Ser Asn Met Ile Gly Ser Ser Pro Ala Asp Gly Leu  
 20 25 30

Leu Ala Leu Pro Leu Leu Thr Val Asp Leu Ala Val Tyr Arg Gly Asn  
 35 40 45

Arg Asp Arg Phe Leu Ala Leu Val Ser Ala His Gly Ala Lys Ala Ala  
 50 55 60

25 Pro His Ala Lys Thr Pro Met Cys Pro Glu Ile Ala Ile Asp Leu Ile  
 65 70 75 80

Glu Ala Gly Ala Trp Gly Ala Thr Val Ala Asp Leu Phe Gln Ala Glu  
 85 90 95

30 Val Leu Leu Lys Ala Gly Val Ser Asn Ile Leu Ile Ala Asn Gln Ile  
 100 105 110

Gly Gly Leu Thr Ser Ala Arg Arg Leu Arg Met Leu Ala Asp Ala Phe  
 115 120 125

35 Pro Lys Ala Glu Ile Ile Cys Cys Val Asp Ser Val Gln Ala Ser Ala  
 130 135 140

40 Asn Leu Val Gln Ala Phe Gln Gly Arg Val Asp Ala Pro Phe Lys Val  
 145 150 155 160

Phe Ile Glu Val Gly Val Gly Arg Thr Gly Ala Arg Thr Leu Asn Val  
 165 170 175

45 Ala Lys Asp Ile Ile Asp Thr Ile Ser Thr Ser Ala Glu Ile Val Leu  
 180 185 190

Ala Gly Val Ser Thr Tyr Glu Gly Ser Val Ser Gly Glu Thr Ser Glu  
 195 200 205

50 Ala Leu Asp Ala Asn Met Ala Ala Leu Phe Asp Leu Leu Thr Asp Ser  
 210 215 220

55

# EP 1 491 549 A1

Leu Ala Ser Ile Arg Glu Lys Asp Pro Gly Arg Pro Leu Thr Val Ser  
 225 230 235 240  
 5  
 Ala Gly Gly Ser Ile His Phe Asp Arg Val Leu Ala Ala Leu Val Pro  
 245 250 255  
 Val Cys Glu Ala Asp Gly Asn Ala Thr Leu Leu Leu Arg Ser Gly Ala  
 260 265 270  
 10  
 Ile Phe Phe Ser Asp His Gly Val Tyr Gln Arg Gly Phe Gln Ala Val  
 275 280 285  
 Asp Ala Arg Asn Leu Leu Ala Ser Gly Lys Val Val Phe Lys Ala Ser  
 290 295 300  
 15  
 Glu Ala Phe Gln Pro Ser Met Arg Ile Trp Ala Glu Val Ile Ser Val  
 305 310 315 320  
 20  
 Pro Glu Pro Gly Leu Ala Ile Val Gly Met Gly Met Arg Asp Val Ser  
 325 330 335  
 Phe Asp Gln Asp Leu Pro Val Ala Leu Arg Leu His Arg Asp Gly His  
 340 345 350  
 25  
 Leu Val Glu Ala Asp Leu Ser Ser Ser Ala Lys Val Gly Lys Leu Asn  
 355 360 365  
 Asp Gln His Ala Phe Leu Ser Phe Gly Asn Gly Ser Ser Leu Ala Ile  
 370 375 380  
 30  
 Gly Asp Val Ile Glu Phe Gly Ile Ser His Pro Cys Thr Cys Phe Asp  
 385 390 395 400  
 35  
 Arg Trp Arg Val Phe His Gly Ile Asp Gly Ser Gly Arg Ile Gln Arg  
 405 410 415  
 Ile Tyr Thr Thr Phe Phe His  
 420  
 40  
 <210> 5  
 <211> 32  
 <212> DNA  
 45  
 <213> artificial sequence  
 <220>  
 50  
 <223> oligonucleotide primer  
 <400> 5  
 cccttaatta atgacgacat ctgataatct tc

32

5           <210> 6  
           <211> 30  
           <212> DNA  
           <213> artificial sequence  
  
 10           <220>  
           <223> oligonucleotide primer  
           <400> 6  
 15           tttgcgggcgg cttagtgggtt atcgcgcgggc 30  
  
           <210> 7  
           <211> 31  
           <212> DNA  
 20           <213> artificial sequence  
  
           <220>  
 25           <223> oligonucleotide primer  
           <400> 7  
           cccgggtacca tgacgacatc tgataatctt c 31  
  
 30           <210> 8  
           <211> 32  
           <212> DNA  
           <213> artificial sequence  
 35           <220>  
           <223> oligonucleotide primer  
 40           <400> 8  
           cccttaatta atgcagtctt cttcagctct tc 32  
  
           <210> 9  
 45           <211> 35  
           <212> DNA  
           <213> artificial sequence  
  
 50           <220>  
           <223> oligonucleotide primer  
           <400> 9  
 55

tttgcggccg cctagtgaaa gaaggttgtg tagat

35

5 <210> 10

<211> 34

<212> DNA

10 <213> artificial sequence

<220>

15 <223> oligonucleotide primer

<400> 10

aaatcatgac tatgcagtct tcttcagctc ttcg

34

20 <210> 11

<211> 33

<212> DNA

25 <213> artificial sequence

<220>

30 <223> oligonucleotide primer

<400> 11

tatagatctc tagtgaaaga aggttggtga gat

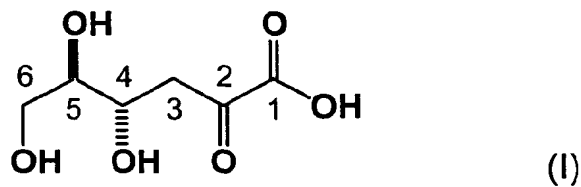
33

35

## Claims

- 40 1. A method for producing 2'-deoxynucleosides or 2'-deoxynucleoside precursors from a compound of formula (I) or its salts

45

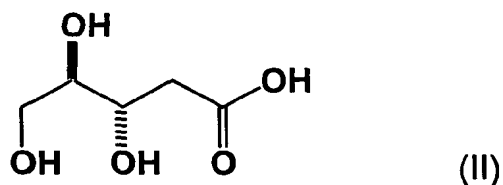


50

or a protected form thereof in a process comprising a decarboxylation step.

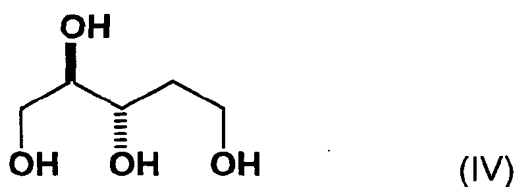
- 55 2. The method of claim 1 wherein the decarboxylation step cleaves the C1-C2 bond of the compound of formula (I) or its salts or a protected form thereof.
3. The method of claim 1 or 2, wherein the decarboxylation step is directly carried out on the compound of formula (I) or its salts or a protected form thereof.

4. The method of any of claims 1 to 3, wherein the decarboxylation step takes place by reacting the compound of formula (I) or its salts or a protected form thereof with hydrogen peroxide to yield a compound of formula (II) or its salts



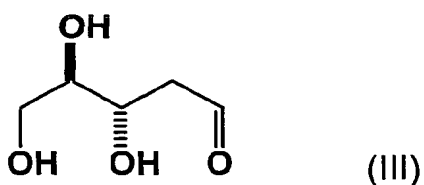
or a protected form thereof as a 2'-deoxynucleoside precursor.

5. The method of claim 4, further comprising the conversion of the compound of formula (II) or its salts or a protected form thereof into a compound of formula (IV)



or a protected form thereof as a 2'-deoxynucleoside precursor.

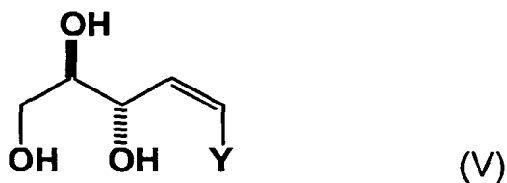
6. The method of claim 4, further comprising the conversion of the compound of formula (II) or its salts or a protected form thereof into a compound of formula (III)



or a protected form thereof as a 2'-deoxynucleoside precursor.

7. The method of claim 6, comprising the conversion of the compound of formula (II) or its salts or a protected form thereof into the compound of formula (IV) or a protected form thereof as an intermediate which is then converted to the compound of formula (III) or a protected form thereof.

8. The method of any of claims 1 to 3, wherein the decarboxylation step takes place by reacting the compound of formula (I) or its salts or a protected form thereof with an amine Y-H, wherein H represents a hydrogen atom bound to the nitrogen atom of the amino group, to produce a compound of formula (V),

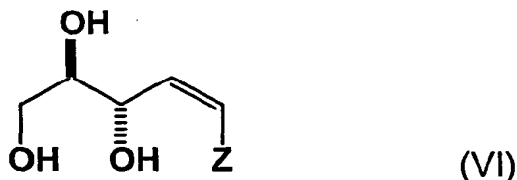


10 or its respective trans isomer or a protected form thereof, as a 2'-deoxynucleoside precursor.

9. The method of claim 8, wherein Y-H represents a linear or cyclic secondary amine.

10. The method of claims 8 or 9, wherein Y-H is morpholine, pyrrolidine, piperidine, N-methyl piperazine or diethyl-  
15 amine.

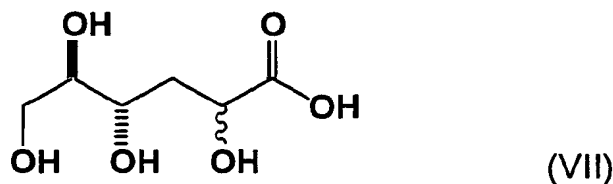
11. The method of any of claims 8 to 10, further comprising the step of reacting a compound of formula (V) or its trans  
20 isomer or a protected form thereof with Z-H, wherein H represents a hydrogen atom and Z represents a leaving group, to produce a compound of formula (VI)



30 or its respective trans isomer or a protected form thereof, as a 2'-deoxynucleoside precursor.

12. The method of claim 11, wherein Z-H is water, to produce a compound of formula (III) or a protected form thereof  
35 as a 2'-deoxynucleoside precursor.

13. The method of claim 1 or 2, wherein the compound of formula (I) or its salts or a protected form thereof is converted  
40 to a compound of formula (VII), or its salts or a protected form thereof or a mixture of the respective epimers,

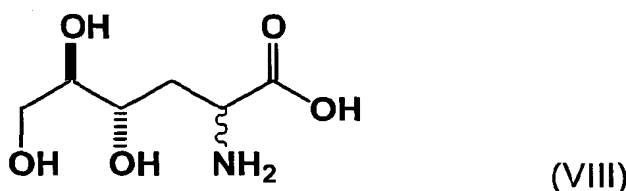


50 which is then decarboxylated to yield a compound of formula (III) or a protected form thereof as a 2'-deoxynucleoside precursor.

14. The method of claim 13, wherein the conversion of (I) or its salts or a protected form thereof to (VII) or a protected  
55 form thereof takes place by reduction with sodium borohydride.

15. The method of claim 13 to 14, wherein the decarboxylation step takes place by reaction with hydrogen peroxide.

16. The method of claim 1 or 2, wherein the compound of formula (I) or its salts or a protected form thereof is converted to a compound of formula (VIII), or its salts or a protected form thereof or a mixture of the respective epimers,



- 15 which is then decarboxylated to yield a compound of formula (III) or a protected form thereof as a 2'-deoxynucleoside precursor.
17. The method of claim 16, wherein a compound of formula (VIII) or a protected form thereof or a mixture of the respective epimers is reacted with ninhydrin, thereby leading to the compound (III) or a protected form thereof.
18. The method of claim 16 or 17, wherein the conversion of (I) or its salts or a protected form thereof to (VIII) or a protected form thereof takes place by reductive amination with ammonia and sodium cyanoborohydride.
19. The method of any of claims 1 to 18, wherein the protective group(s) are independently chosen from acetate ester, benzoate ester, allyl ether, benzyl ether, trityl ether, ter-butyldimethylsilyl (TBDMS) ether, isopropylidene or a benzylidene acetal.
20. The method of any one of claims 1 to 19, comprising the preliminary step of producing the compound of formula (I) from a D-gluconate salt by the use of a gluconate dehydratase activity.
21. The method of claim 20, wherein the D-gluconate salt is potassium or sodium D-gluconate.
22. The method of claims 20 or 21, wherein the gluconate dehydratase is encoded by a polynucleotide comprising the nucleotide sequence selected from the group consisting of:
- 35 (a) nucleotide sequences encoding a polypeptide comprising the amino acid sequence of SEQ ID N°2;  
 (b) nucleotide sequences comprising the coding sequence of SEQ ID N°1;  
 (c) nucleotide sequences encoding a fragment encoded by a nucleotide sequence of (a) or (b);  
 (d) nucleotide sequences hybridising with a nucleotide sequence of any one of (a) to (c); and  
 40 (e) nucleotide sequences which deviate from the nucleoside sequence of (d) as a result of degeneracy of the genetic code.
23. The method of any one of claims 1 to 19, comprising the preliminary step of producing the compound of formula (I) from D-glucosamine by the use of a glucosamine deaminase activity.
24. The method of claim 23, wherein the glucosamine deaminase is encoded by a polynucleotide comprising the nucleotide sequence selected from the group consisting of:
- 45 (a) nucleotide sequences encoding a polypeptide comprising the amino acid sequence of SEQ ID N°4;  
 (b) nucleotide sequences comprising the coding sequence of SEQ ID N°3;  
 (c) nucleotide sequences encoding a fragment encoded by a nucleotide sequence of (a) or (b);  
 (d) nucleotide sequences hybridising with a nucleotide sequence of any one of (a) to (c); and  
 50 (e) nucleotide sequences which deviate from the nucleoside sequence of (d) as a result of degeneracy of the genetic code.
25. Use of a polynucleotide as defined in claim 22 or of a gluconate dehydratase encoded by such a polynucleotide in a method according to claims 20 or 21.
- 55

- 26.** Use of a polynucleotide as defined in claim 24 or of a glucosaminase deaminase encoded by such a polynucleotide in a method according to claim 23.

5

10

15

20

25

30

35

40

45

50

55





European Patent  
Office

## EUROPEAN SEARCH REPORT

Application Number  
EP 03 01 3457

DOCUMENTS CONSIDERED TO BE RELEVANT			
Category	Citation of document with indication, where appropriate, of relevant passages	Relevant to claim	CLASSIFICATION OF THE APPLICATION (Int.Cl.7)
A	SHELTON M C ET AL: "2-Keto-3-deoxy-6-phosphogluconate aldolases as catalysts for stereocontrolled carbon-carbon bond formation" JOURNAL OF THE AMERICAN CHEMICAL SOCIETY, AMERICAN CHEMICAL SOCIETY, WASHINGTON, DC, US, vol. 118, no. 9, 6 March 1996 (1996-03-06), pages 2117-2125, XP002263455 ISSN: 0002-7863 Scheme 6	1	C07H7/027 C07H1/00
A	US 5 872 247 A (DUFLOT PIERRICK ET AL) 16 February 1999 (1999-02-16) exemples	1	
A	US 5 846 794 A (DELOBEAU DIDIER ET AL) 8 December 1998 (1998-12-08) exemple 3	1	
A	H. KILIANI, H. NAEGEL: "Ueber Meta- und Parasaccharin" CHEM. BER., vol. 35, 1902, pages 3528-3533, XP002267849 * page 3531 - page 3532 *	13	TECHNICAL FIELDS SEARCHED (Int.Cl.7) C07H
The present search report has been drawn up for all claims			
Place of search THE HAGUE		Date of completion of the search 23 January 2004	Examiner de Nooy, A
CATEGORY OF CITED DOCUMENTS X : particularly relevant if taken alone Y : particularly relevant if combined with another document of the same category A : technological background O : non-written disclosure P : intermediate document		T : theory or principle underlying the invention E : earlier patent document, but published on, or after the filing date D : document cited in the application L : document cited for other reasons & : member of the same patent family, corresponding document	

EPO FORM 1503 03 82 (P04C01)

**ANNEX TO THE EUROPEAN SEARCH REPORT  
ON EUROPEAN PATENT APPLICATION NO.**

EP 03 01 3457

This annex lists the patent family members relating to the patent documents cited in the above-mentioned European search report.  
The members are as contained in the European Patent Office EDP file on  
The European Patent Office is in no way liable for these particulars which are merely given for the purpose of information.

23-01-2004

Patent document cited in search report		Publication date		Patent family member(s)	Publication date
US 5872247	A	16-02-1999	FR	2749306 A1	05-12-1997
			AT	203996 T	15-08-2001
			CA	2206390 A1	03-12-1997
			DE	69705995 D1	13-09-2001
			DE	69705995 T2	04-04-2002
			DK	811632 T3	12-11-2001
			EP	0811632 A1	10-12-1997
			ES	2162211 T3	16-12-2001
			JP	10081693 A	31-03-1998
-----					
US 5846794	A	08-12-1998	FR	2749307 A1	05-12-1997
			AT	212383 T	15-02-2002
			CA	2206389 A1	29-11-1997
			DE	69709985 D1	14-03-2002
			EP	0810292 A1	03-12-1997
			JP	10087531 A	07-04-1998
-----					